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Reviewer: Durreshwar Anjum
Timestamp: Fri Oct 19 11:38:00 EDT 2007
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* *
Reviewer Comments:
- 6 -

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Application No: 09426011   Version No: 6.0

Input Set:

Output Set:

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<110>   Simons, Michael
        Gao, Youhe

<120>   Method for PR-39 Peptide Regulated Stimulation of Angiogenesis

<130>   BI-0004US.P1

<140>   09426011
<141>   1999-10-25

<150>   US 09/276,868
<151>   1999-03-26

<160>   16

<170>   PatentIn version 3.3

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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg Pro Arg Pro Pro Pro
1   5   10   15

Phe Phe Pro Pro Arg Leu Pro Pro Arg Ile Pro Pro Gly Phe Pro Pro
            20   25   30

Arg Phe Pro Pro Arg Phe Pro
        35

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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg Pro Arg Pro Pro Pro
1   5   10   15

Phe Phe Pro Pro Arg Leu Pro Pro Arg Ile Pro Pro Gly Phe Pro Pro

Arg Phe Pro Pro Arg Phe Pro
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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg Pro Arg Pro Pro
1   5   10   15

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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg
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Arg Arg Arg Pro Arg Pro Pro Tyr
1   5

<210>   6
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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg Pro Arg Pro Pro Pro
1   5   10   15

Phe Phe Pro Pro Arg Leu Pro Pro Arg Ile Pro Pro Gly Phe Pro Pro
            20   25   30

Arg Phe Pro Pro Arg Phe Pro
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Arg Arg Arg Pro Arg Pro Pro Tyr Leu Pro Arg
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Met Ser Ser Ile Gly Thr Gly Tyr Asp Leu Ser Ala Ser Thr Phe Ser
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Pro Asp Gly Arg Val Phe Gln Val Glu Tyr Ala Met Lys Ala Val Glu
            20   25   30

Asn Ser Ser Thr Ala Ile Gly Ile Arg Cys Lys Asp Gly Val Val Phe
        35   40   45

Gly Val Glu Lys Leu Val Leu Ser Lys Leu Tyr Glu Glu Gly Ser Asn
    50   55   60

Lys Arg Leu Phe Asn Val Asp Arg His Val Gly Met Ala Val Ala Gly
65   70   75   80

Leu Leu Ala Asp Ala Arg Ser Leu Ala Asp Ile Ala Arg Glu Glu Ala
                85   90   95

Ser Asn Phe Arg Ser Asn Phe Gly Tyr Asn Ile Pro Leu Lys His Leu
            100   105   110

Ala Asp Arg Val Ala Met Tyr Val His Ala Tyr Thr Leu Tyr Ser Ala
            115   120   125

Val Arg Pro Phe Gly Cys Ser Phe Met Leu Gly Ser Tyr Ser Ala Asn
        130   135   140

Asp Gly Ala Gln Leu Tyr Met Ile Asp Met Ser Gly Val Ser Tyr Gly
145   150   155   160

Tyr Trp Gly Cys Ala Ile Gly Lys Ala Arg Gln Ala Ala Lys Thr Glu
            165   170   175

Ile Glu Lys Leu Gln Met Lys Glu Met Thr Cys Arg Asp Val Val Lys
            180   185   190

Glu Val Ala Lys Ile Ile Tyr Ile Val His Asp Glu Val Lys Asp Lys
            195   200   205

Ala Phe Glu Leu Glu Leu Ser Trp Val Gly Glu Leu Thr Lys Gly Arg
        210   215   220

His Glu Ile Val Pro Lys Asp Ile Arg Glu Glu Ala Glu Lys Tyr Ala
225   230   235   240

Lys Glu Ser Leu Lys Glu Glu Asp Glu Ser Asp Asp Asp Asn Met
            245   250   255

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Ala
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Glu Lys Glu Lys Glu Glu Asn Glu Lys Lys Lys Gln Lys Lys Ala Ser
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Pro Ala Glu Lys Ala Asp Glu Pro Met Glu His
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Ala Lys Glu Ser Leu Lys Glu Glu Asp Glu Ser Asp Asp Asp Asn Met
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Pro Pro Xaa Xaa Pro Pro Xaa Xaa Pro
1   5

<210> 16
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<223> Xaa denotes any amino acid residue

<400> 16

Pro Pro Xaa Xaa Xaa Pro Pro Xaa Xaa Pro
1   5   10